RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10

Source:

Date Processed by STIC:

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IFW16

RAW SEQUENCE LISTING DATE: 11/29/2006
PATENT APPLICATION: US/10/619,939A TIME: 13:49:13

Input Set : R:\Revised Sequence Listing 11-21-06.ST25.txt

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3 <110> APPLICANT: Advisys, Inc.
     5 <120> TITLE OF INVENTION: Codon optimized Synthetic Plasmid
     7 <130> FILE REFERENCE: 108328.00146
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/619,939A
C--> 9 <141> CURRENT FILING DATE: 2003-07-15
     9 <160> NUMBER OF SEQ ID NOS: 43
    11 <170> SOFTWARE: PatentIn version 3.3
    13 <210> SEQ ID NO: 1
    14 <211> LENGTH: 3534
    15 <212> TYPE: DNA
    16 <213> ORGANISM: artificial sequence
    18 <220> FEATURE:
    19 <223> OTHER INFORMATION: Plasmid vector having an analog GHRH sequence.
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    24 accgcggtgg cggccgtccg ccctcggcac catcctcacg acacccaaat atggcgacgg
                                                                            120
    26 qtqaqqaatq qtqqqqaqtt atttttaqaq cggtqaggaa ggtgggcagg cagcaggtgt
                                                                            180
    28 tggcgctcta aaaataactc ccgggagtta tttttagagc ggaggaatgg tggacaccca
                                                                            240
    30 aatatggcga cggttcctca cccgtcgcca tatttgggtg tccgccctcg gccggggccg
                                                                            300
    32 catteetggg ggeeggegg tgeteeegee egeetegata aaaggeteeg gggeeggegg
                                                                            360
     34 cqqcccacqa qctacccqqa qqaqcqqqaq gcqccaagct ctagaactag tggatcccaa
                                                                            420
    36 ggcccaactc cccgaaccac tcagggtcct gtggacagct cacctagctg ccatggtgct
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    38 ctgggtgttc ttctttgtga tcctcaccct cagcaacage tcccactgct ccccacctcc
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     40 ccctttgacc ctcaggatgc ggcggcacgt agatgccatc ttcaccaaca gctaccggaa
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                                                                            660
     42 qqtqctqqcc caqctgtccq cccqcaaqct gctccaggac atcctgaaca ggcagcaggg
                                                                            720
     44 agaqaqqaac caagagcaag gagcataatg actgcaggaa ttcgatatca agcttatcgg
                                                                            780
     46 ggtggcatcc ctgtgacccc tccccagtgc ctctcctggc cctggaagtt gccactccag
     48 tycccaccag ccttytccta ataaaattaa gttycatcat tttytctyac tagytytcct
                                                                            840
     50 tctataatat tatggggtgg aggggggtgg tatggagcaa ggggcaagtt gggaagacaa
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     52 cctgtagggc ctgcggggtc tattgggaac caagctggag tgcagtggca caatcttggc
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     54 tractgraat ctrcgcctrc tgggttraag cgattrtctct gertragert crcgagttgt
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     56 tgggattcca ggcatgcatg accaggctca gctaattttt gtttttttgg tagagacggg
                                                                           1080
                                                                           1140
     58 gtttcaccat attggccagg ctggtctcca actcctaatc tcaggtgatc tacccacctt
     1200
     62 ttttaaaata actataccag caggaggacg tccagacaca gcataggcta cctggccatg
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     64 cccaaccggt gggacatttg agttgcttgc ttggcactgt cctctcatgc gttgggtcca
     66 ctcagtagat gcctgttgaa ttcgataccg tcgacctcga gggggggccc ggtaccagct
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     68 tttgttccct ttagtgaggg ttaatttcga gcttggcgta atcatggtca tagctgtttc
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     70 ctgtgtgaaa ttgttatccg ctcacaattc cacacaacat acgagccgga agcataaagt
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     72 gtaaagcctg gggtgcctaa tgagtgagct aactcacatt aattgcgttg cgctcactgc
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     74 ccgctttcca gtcgggaaac ctgtcgtgcc agctgcatta atgaatcggc caacgcgcqg
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     76 qqaqaqqqqq tttqcqtatt qgqcqctctt ccqcttcctc qctcactqac tcqctqcqct
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     78 cqqtcqttcq gctgcggcga qcggtatcag ctcactcaaa ggcggtaata cggttatcca
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Input Set : E:\Revised Sequence Listing 11-21-06.ST25.txt

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                                                                        1860
84 acaaaaatcg acgctcaagt cagaggtggc gaaacccgac aggactataa agataccagg
                                                                        1920
86 cgtttccccc tggaagctcc ctcgtgcgct ctcctgttcc gaccctgccg cttaccggat
                                                                        1980
88 acctgtccgc ctttctccct tcgggaagcg tggcgctttc tcatagctca cgctgtaggt
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90 atctcagttc ggtgtaggtc gttcgctcca agctgggctg tgtgcacgaa ccccccgttc
                                                                        2100
92 agogogacco otgogootta tooggtaact atogtottga gtocaacagg gtaagacacg
94 acttategee actggeagea gecaetggta acaggattag cagagegagg tatgtaggeg 2220
96 gtgctacaga gttcttgaag tggtggccta actacggcta cactagaaga acagtatttg
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98 qtatctqcqc tctqctqaaq ccaqttacct tcggaaaaag agttggtagc tcttgatccg
100 gcaaacaaac caccgctggt agcggtggtt tttttgtttg caagcagcag attacgcgca
                                                                         2400
                                                                         2460
102 gaaaaaaagg atctcaagaa gatcctttga tcttttctac ggggtctgac gctcagaaga
104 actcqtcaaq aaqqcqataq aaqqcqatgc gctgcgaatc gggagcggcg ataccgtaaa
                                                                         2520
106 gcacqaqqaa gcqqtcaqcc cattcqccqc caagctcttc agcaatatca cgggtagcca
                                                                         2580
108 acgctatgtc ctgatagcgg tccgccacac ccagccggcc acagtcgatg aatccagaaa
                                                                         2640
                                                                         2700
110 aggggggatt ttccaccatq atattcggca aggaggcatc gccatgggtc acgacgagat
112 cetequeqte gggcatgeqe geettqaqee tggcgaacag tteggetgge gegageeeet
                                                                         2760
114 gatgetette gtecagatea teetgatega caagacegge ttecateega gtacgtgete
                                                                         2820
116. gobogatgog atgittteget tiggtggtega atgggeaggt ageeggatea agegtatigea
                                                                         2830
                                                                         2940
118 geogeogeat tgeateagee atgatggata etttetegge aggageaagg tgagatgaea
120 qqaqatcctq cccqqcact tcqcccaata gcaqccaqtc ccttcccqct tcaqtqacaa
                                                                         3000
122 cgtcgagcac agctgcgcaa ggaacgcccg tcgtggccag ccacgatagc cgcgctgcct
                                                                         3060
124 cgtcctgcag ttcattcagg gcaccggaca ggtcggtctt gacaaaaaga accgggcgcc
                                                                         3120
126 cctgcgctga cagccggaac acggcggcat cagagcagcc gattgtctgt tgtgcccagt
                                                                         3180
128 catagoogaa tagoototoo acccaagogg coggagaaco tgogtgcaat coatottgtt
                                                                         3240
130 caatcatgcg aaacgatcct catcctgtct cttgatcaga tcttgatccc ctgcgccatc
                                                                         3300
132 agateettgg eggeaagaaa gecateeagt ttaetttgea gggetteeea acettaeeag
                                                                         3360
134 agggegeece agetggeaat teeggttege ttgetgteea taaaacegee cagtetagea
                                                                         3420
                                                                         3480
136 actgttggga agggcgatcg gtgcgggcct cttcgctatt acgccagctg gcgaaagggg
138 gatqtgctqc aaggcgatta agttgggtaa cgccagggtt ttcccagtca cgac
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141 <210> SEQ ID NO: 2
142 <211> LENGTH: 2739
143 <212> TYPE: DNA
144 <213> ORGANISM: artificial sequence
146 <220> FEATURE:
147 <223> OTHER INFORMATION: Optimized vector having an analog GHRH sequence.
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152 gggtgaggaa tggtggggag ttatttttag agcggtgagg aaggtgggca ggcagcaggt
                                                                          120
154 gttggcgctc taaaaataac tcccgggagt tatttttaga gcggaggaat ggtggacacc
                                                                          180
156 caaatatggc gacggtteet caccegtege catatttggg tgtccgccct cggccggggc
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158 cgcattcctg ggggccgggc ggtgctcccg cccgcctcga taaaaggctc cggggccggc
                                                                          300
                                                                          360
160 ggcggcccac gagctacccg gaggagcggg aggcgccaag cggatcccaa ggcccaactc
162 cccgaaccac tcagggtcct gtggacagct cacctagctg ccatggtgct ctgggtgttc
                                                                          420
164 ttctttgtga tcctcaccct cagcaacagc tcccactgct ccccacctcc ccctttgacc
                                                                          480
166 ctcaggatgc ggcggtatgc agatgccatc ttcaccaaca gctaccggaa ggtgctgggc
                                                                          540
168 cagctgtccg cccgcaagct gctccaggac atcatgagca ggcagcaggg agagaggaac
                                                                          600
170 caagagcaag gagcataatg actgcaggaa ttcgatatca agcttatcgg ggtggcatcc
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172 ctgtgacccc tccccagtgc ctctcctggc cctggaagtt gccactccag tgcccaccag
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Input Set : E:\Revised Sequence Listing 11-21-06.ST25.txt

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174 ccttgtccta ataaaattaa gttgcatcat tttgtctgac taggtgtcct tctataatat
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 176 tatggggtgg agggggtgg tatggagcaa ggggcaagtt gggaagacaa cctgtagggc
 178 tcgaggggg gcccggtacc agcttttgtt ccctttagtg agggttaatt tcgagcttgg
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                                                                           960
 180 tetteegett cetegeteac tgactegetg egeteggteg tteggetgeg gegageggta
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 182 tcagctcact caaaggcggt aatacggtta tccacagaat caggggataa cgcaggaaag
                                                                          1080
 184 aacatgtgag caaaaggcca gcaaaaggcc aggaaccgta aaaaggccgc gttgctggcg
 186 tttttccata ggctccgccc ccctgacgaq catcacaaaa atcgacgctc aagtcagagg
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 188 tggcgaaacc cgacaggact ataaagatac caggcgtttc cccctggaag ctccctcgtg
                                                                          1200
                                                                          1260
 190 egeteteetg tteegaeeet geegettaee ggataeetgt eegeetttet eeetteggga
                                                                          1320
 192 agcgtggcgc tttctcatag ctcacgctgt aggtatctca gttcggtgta ggtcgttcgc
 194 tocaagetgg getgtgtgca egaaceeece gtteageeeg acegetgege ettateeggt
                                                                          1380
                                                                          1440
 196 aactatcgtc ttgagtccaa cccggtaaga cacgacttat cgccactggc agcagccact
                                                                          1500
 198 qqtaacagga ttagcagagc gaggtatgta ggcggtgcta cagagttctt gaagtggtgg
                                                                          1560
 200 cctaactacg gctacactag aagaacagta tttggtatct gcgctctgct gaagccagtt
 202 accttcqqaa aaagagttgg tagctcttga tccggcaaac aaaccaccgc tggtagcggt
                                                                          1620
                                                                          1680
 204 gqtttttttg tttgcaagca gcagattacg cgcagaaaaa aaggatctca agaagatcct
 206 ttgatctttt ctacggggtc tgacgctcag ctagcgctca gaagaactcg tcaagaaggc
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                                                                          1800
 208 gatagaaggc gatgcgctgc gaatcgggag cggcgatacc gtaaagcacg aggaagcggt
                                                                          1860
210 cagoccatto geogecaago tettoagoaa tatgaegggt agecaaeget atgtoetgat
                                                                                 1 144- 1
                                                                          1920
 212 agoggtocgo cacaccoago oggocacagt ogatgaatoo agaaaagogg coattitoca
 214 ccatgatatt cggcaagcag gcatcgccat gagtcacgac gagatcctcg ccgtcgggca
                                                                          1980
 216 tgcgcgcctt gagcctggcg aacagttcgg ctggcgcgag cccctgatgc tcttcgtcca
                                                                          2040
 218 gatcatectg ategacaaga eeggetteea teegagtaeg tgetegeteg atgegatgtt
                                                                          2100
                                                                          2160
 220 tegettggtg gtegaatggg caggtageeg gateaagegt atgeageege egeattgeat
 222 cagccatgat ggatactttc tcggcaggag caaggtgaga tgacaggaga tcctgccccg
                                                                          2220
                                                                          2280
 224 gcacttegee caatageage cagtecette eegetteagt gacaaegteg ageacagetg
                                                                          2340
 226 cgcaaggaac gcccgtcgtg gccagccacg atagccgcgc tgcctcgtcc tgcagttcat
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 230 ggaacacggc ggcatcagag cagccgattg tctgttgtgc ccagtcatag ccgaatagcc
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 232 totocaccca agoggoogga gaacctgogt gcaatccatc ttgttcaatc atgcgaaacg
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 234 atcctcatcc tgtctcttga tcagatcttg atcccctgcg ccatcagatc cttggcggca
                                                                          2580
                                                                          2640
 236 aqaaaqccat ccaqtttact ttqcagggct tcccaacctt accagagggc gccccagctg
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 238 qcaattccqq ttcqcttqct gtccataaaa ccgcccagtc tagcaactgt tgggaagggc
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 240 gatcgtgtaa tacgactcac tatagggcga attggagct
 243 <210> SEQ ID NO: 3
 244 <211> LENGTH: 795
 245 <212> TYPE: DNA
 246 <213> ORGANISM: artificial sequence
 248 <220> FEATURE:
 249 <223> OTHER INFORMATION: Nucleic acid sequence for the antibiotic resistance gene
           kanamycin.
 250
 252 <400> SEQUENCE: 3
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 255 ggctatgact gggcacaaca gacaatcggc tgctctgatg ccgccgtgtt ccggctgtca
                                                                            120
 257 gegeagggge geoeggttet ttttgteaag accgaectgt eeggtgeect gaatgaactg
                                                                            180
 259 caggacgagg cagcgcggct atcgtggctg gccacgacgg gcgttccttg cgcagctgtg
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                                                                           300
 261 ctcgacgttg tcactgaagc gggaagggac tggctgctat tgggcgaagt gccggggcag
 263 gatctcctgt catctcacct tgctcctgcc gagaaagtat ccatcatggc tgatgcaatg
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Input Set : E:\Revised Sequence Listing 11-21-06.ST25.txt

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    271 ggcgaggate tegtegtgae teatggegat geetgettge egaatateat ggtggaaaat
                                                                               600
    273 ggccgctttt ctggattcat cgactgtggc cggctgggtg tggcggaccg ctatcaggac
                                                                               660
                                                                               720
    275 atagcgttgg ctacccgtga tattgctgaa gagcttggcg gcgaatgggc tgaccgcttc
    277 ctcgtgcttt acggtatcgc cgctcccgat tcgcagcgca tcgccttcta tcgccttctt
                                                                               780
2. 279 gacgagttct tctga
                                                                               795
                                            A representation of the second
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    283 <211> LENGTH: 219
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    285 <213> ORGANISM: artificial sequence
    287 <220> FEATURE:
    288 <223> OTHER INFORMATION: Sequence for an analog porcine GHRH sequence.
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    295 taccqqaagg tgctggccca gctgtccgcc cgcaagctgc tccaggacat cctgaacagg
                                                                               180
                                                                               219
    297 cagcagggag agaggaacca agagcaagga gcataatga
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                                                      - 2:
    301 <211> LENGTH: 246
    302 <212> TYPE: DNA
    303 <213> ORGANISM: artificial sequence
    305 <220> FEATURE:
    306 <223> OTHER INFORMATION: Sequence for an analog mouse GHRH sequence.
     308 <400> SEQUENCE: 5
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    311 ctgcctccca gccctccctt caggatgcag aggcacgtgg acgccatctt caccaccaac
                                                                               120
                                                                               180
    313 tacaggaagc tgctgagcca gctgtacgcc aggaaggtga tccaggacat catgaacaag
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    315 cagggcgaga ggatccagga gcagagggcc aggctgagct gataagcttg cgatgagttc
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    317 ttctaa
    320 <210> SEQ ID NO: 6
     321 <211> LENGTH: 234
     322 <212> TYPE: DNA
    323 <213> ORGANISM: artificial sequence
    325 <220> FEATURE:
    326 <223> OTHER INFORMATION: Sequence for an analog rat GHRH sequence.
    328 <400> SEQUENCE: 6
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     331 ctgcctccca gccctccctt cagggtgcgc cggcacgccg acgccatctt caccagcagc
                                                                               120
     333 tacaqqaqqa teetgqgeca getgtaeget aggaagetee tgeaegagat catgaacagg
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     339 <211> LENGTH: 225
     340 <212> TYPE: DNA
     341 <213> ORGANISM: artificial sequence
     343 <220> FEATURE:
     344 <223> OTHER INFORMATION: Sequence for an analog bovine GHRH sequence.
     346 <400> SEQUENCE: 7
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Input Set : E:\Revised Sequence Listing 11-21-06.ST25.txt

Output Set: N:\CRF4\11292006\J619939A.raw

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426 <211> LENGTH: 782 427 <212> TYPE: DNA VERIFICATION SUMMARY

DATE: 11/29/2006

the second secon

PATENT APPLICATION: US/10/619,939A

TIME: 13:49:14

Input Set : E:\Revised Sequence Listing 11-21-06.ST25.txt

Output Set: N:\CRF4\11292006\J619939A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date